

SEQUENCE LISTING

<110> Currie, Mark G. Mahajan-Miklos, Shalina

<120> METHODS AND COMPOSITIONS FOR THE TREATMENT OF GASTROINTESTINAL DISORDERS

<130> 14184-043001

<140> US 10/796,719

<141> 2004-03-09

<150> US 10/766,735

<151> 2004-01-28

<150> US 60/443,098

<151> 2003-01-28

<150> US 60/471,288

<151> 2003-05-15

<150> US 60/519,460

<151> 2003-11-12

<160> 149

<170> FastSEQ for Windows Version 4.0

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Cys Asn
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Cys Tyr
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Gln Ala Cys Asp Pro Pro Ser Pro Pro Ala Glu Val Ser Ser Asp Trp
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Gln Glu Thr Ala Ser Gly Gln Val Gly Asp Val Ser Ser Ser Thr Ile
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Ala Thr Glu Val Ser Glu Ala Glu Cys Gly Thr Gln Ser Ala Thr Thr
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Ala Cys Phe Gly Cys
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Ser Asp Trp Cys Cys Glu Val Cys Cys Asn Pro Ala Cys Ala Gly Cys
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Ile Asp Cys Cys Glu Ile Cys Cys Asn Pro Ala Cys Phe Gly Cys Leu
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Asn
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Asn
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Cys Tyr
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Ser Phe Ser Gln Ser Thr Glu Ser Leu Asp Ser Ser Lys Glu Lys Ile
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                                25
Thr Leu Glu Thr Lys Lys Cys Asp Val Val Lys Asn Asn Ser Glu Lys
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Lys Ser Glu Asn Met Asn Asn Thr Phe Tyr Cys Cys Glu Leu Cys Cys
Asn Pro Ala Cys Ala Gly Cys Tyr
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Pro Phe Ala Gln Asp Ala Lys Pro Val Glu Ser Ser Lys Glu Lys Ile
Thr Leu Glu Ser Lys Lys Cys Asn Ile Ala Lys Lys Ser Asn Lys Ser
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Gly Pro Glu Ser Met Asn Ser Ser Asn Tyr Cys Cys Glu Leu Cys Cys
Asn Pro Ala Cys Thr Gly Cys Tyr
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Ala Phe Gly Gln Glu Thr Val Ser Gly Gln Phe Ser Asp Ala Leu Ser
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Thr Pro Ile Thr Ala Glu Val Tyr Lys Gln Ala Cys Asp Pro Pro Leu
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Gly Pro Glu Ser Met Asn
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Gly Pro Glu Ser Met
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Thr Leu Glu Ser Lys Lys Cys Asn Ile Val Lys Lys Asn Asn Glu Ser
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Ser Pro Glu Ser Met
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Gly Cys Tyr
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Gly Cys Tyr
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Gly Cys Tyr
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Cys Cys Glu Leu Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
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Cys Cys Glu Tyr Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
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Gly Cys Tyr Asp Phe
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Gly Cys Tyr Asp Phe
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Gly Cys Tyr Asp Phe
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Gly Cys Tyr Asp Phe
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Cys Cys Glu Lys Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr Asp Phe
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Phe
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Phe
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Phe
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cacaccatat gaagaaatca atattattta tttttctttc tg
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cacacctcga gttaggtctc catgctttca ggaccacttt tattac	46
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<210> 66 <211> 21 <212> PRT <213> Artificial Sequence	
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     other than Leu; or Xaa = Phe, Trp, and Tyr; or
      selected from from any other natural or
      non-natural aromatic amino acid; or Xaa = Tyr
<220>
<221>VARIANT
<222> 1, 2, 3, 4, 5
<223> Xaa1 = Asn, Xaa2 = Ser, Xaa3 = Ser, Xaa4 = Asn,
     Xaa5 = Tyr; or Xaa1-Xaa5 is missing; or Xaa1-Xaa4
      is missing; or Xaal -Xaa3 is missing; or Xaal and
     Xaa2 is missing; or Xaal is missing
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<221>VARIANT
<222> 19, 20, 21
<223> Xaa 20 = Asp, Xaa21 = Phe or missing; or Xaa20 =
      Asn or Glu and Xaa21 is missing; or X19-Xaa21 is
      missing
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Xaa Xaa Xaa Xaa Cys Cys Glu Xaa Cys Cys Asn Pro Ala Cys Thr
Gly Cys Tyr Xaa Xaa
            20
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Gly Cys Tyr
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Gly Cys Tyr
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Gly Cys Tyr
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Gly Cys Tyr
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Gly Cys Tyr
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Ser Ser Asn Tyr Cys Cys Glu Tyr Cys Cys Asn Pro Ala Cys Thr Gly
Cys Tyr
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Gly Cys Tyr
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<210> 77

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Gly Cys Tyr
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Gly Cys Tyr
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Gly Cys Tyr
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                 5
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                 5
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Gly Cys Tyr
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Cys Cys Glu Ile Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
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Cys Cys Glu Met Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
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Cys Cys Glu Phe Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
<210> 106
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Cys Cys Glu Pro Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
<210> 107
<211> 14
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Cys Cys Glu Ser Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
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Cys Cys Glu Thr Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
                                     10
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Cys Cys Glu Trp Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
<210> 110
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Cys Cys Glu Val Cys Cys Asn Pro Ala Cys Thr Gly Cys Tyr
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<400> 111
Gln His Asn Pro Arg
<210> 112
<211> 6
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Val Gln His Asn Pro Arg
<210> 113
<211> 7
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<400> 113
Val Arg Gln His Asn Pro Arg
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Val Arg Gly Gln His Asn Pro Arg
<210> 115
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<400> 115
Val Arg Gly Pro Gln His Asn Pro Arg
<210> 116
<211> 10
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Val Arg Gly Pro Arg Gln His Asn Pro Arg
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<210> 117
<211> 11
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Val Arg Gly Pro Arg Arg Gln His Asn Pro Arg
<210> 118
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<400> 118
Arq Gln His Asn Pro Arg
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<211> 21
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<222> 9
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<220>
<221>VARIANT
<222> 8, 12, 13, 14, 16, 17, 19, 20, 21
<223> Xaa = Ala, Arg, Asn, Asp, Cys, Gln, Glu, Gly, His, Ile, Leu,
Lys, Met, Phe, Pro, Ser, Thr, Trp, Tyr or Val
<400> 119
Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Cys Cys Xaa Xaa Cys Xaa
Xaa Cys Xaa Xaa Xaa
            20
<210> 120
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<213> Artificial Sequence
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      Xaa5 = Tyr or missing; or Xaa1- Xaa4 is missing
      and Xaa5 = Asn
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<222> 9
<223> Xaa = Leu, Ile, Val, Trp, Tyr or Phe
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<222> 16
<223> Xaa = Thr, Ala, or Trp
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<221>VARIANT
<222> 19
<223> Xaa = Trp, Tyr, Or Leu or is missing
<220>
<221>VARIANT
<222> 20, 21
<223> Xaa = Asp, Phe
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Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Cys Cys Asn Pro Ala Cys Xaa
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Gly Cys Xaa Xaa Xaa
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<211> 5
<212> PRT
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Asn Ser Ser Asn Tyr
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<210> 122
<211> 30
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<212> PRT
<213> Yersinia enterocolitica
<400> 122
Gln Ala Cys Asp Pro Pro Leu Pro Pro Ala Glu Val Ser Ser Asp Trp
                               10
Asp Cys Cys Asp Val Cys Cys Asn Pro Ala Cys Ala Gly Cys
<210> 123
<211> 6
<212> PRT
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<223> Synthetically generated peptide
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Lys Lys Lys Lys Lys
<210> 124
<211> 7
<212> PRT
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<400> 124
Asp Lys Lys Lys Lys Lys
               5
<210> 125
<211> 13
<212> PRT
<213> Artificial Sequence
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<400> 125
Cys Cys Glu Tyr Cys Cys Asn Pro Ala Cys Thr Gly Cys
                 5
<210> 126
<211> 13
<212> PRT
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<223> Synthetically generated peptide
<400> 126
Cys Cys Glu Ala Cys Cys Asn Pro Ala Cys Thr Gly Cys
                5
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<211> 13
<212> PRT
<213> Artificial Sequence
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<223> Synthetically generated peptide
<400> 127
Cys Cys Glu Arg Cys Cys Asn Pro Ala Cys Thr Gly Cys
<210> 128
<211> 13
<212> PRT
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<400> 128
Cys Cys Glu Asn Cys Cys Asn Pro Ala Cys Thr Gly Cys
<210> 129
<211> 13
<212> PRT
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Cys Cys Glu Asp Cys Cys Asn Pro Ala Cys Thr Gly Cys
                5
<210> 130
<211> 13
<212> PRT
<213> Artificial Sequence
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<400> 130
Cys Cys Glu Cys Cys Cys Asn Pro Ala Cys Thr Gly Cys
                5
<210> 131
<211> 13
<212> PRT
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<400> 131
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Cys Cys Glu Glu Cys Cys Asn Pro Ala Cys Thr Gly Cys
<210> 133
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Cys Cys Glu Gly Cys Cys Asn Pro Ala Cys Thr Gly Cys
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<211> 13
<212> PRT
<213> Artificial Sequence
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<400> 134
Cys Cys Glu His Cys Cys Asn Pro Ala Cys Thr Gly Cys
                 5
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<400> 135
Cys Cys Glu Ile Cys Cys Asn Pro Ala Cys Thr Gly Cys
                5
<210> 136
<211> 13
<212> PRT
<213> Artificial Sequence
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Cys Cys Glu Lys Cys Cys Asn Pro Ala Cys Thr Gly Cys
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Cys Cys Glu Met Cys Cys Asn Pro Ala Cys Thr Gly Cys
<210> 138
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Cys Cys Glu Phe Cys Cys Asn Pro Ala Cys Thr Gly Cys
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Cys Cys Glu Pro Cys Cys Asn Pro Ala Cys Thr Gly Cys
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<400> 140
Cys Cys Glu Ser Cys Cys Asn Pro Ala Cys Thr Gly Cys
<210> 141
<211> 13
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<223> Synthetically generated peptide
<400> 141
Cys Cys Glu Thr Cys Cys Asn Pro Ala Cys Thr Gly Cys
                5
<210> 142
<211> 13
<212> PRT
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Cys Cys Glu Trp Cys Cys Asn Pro Ala Cys Thr Gly Cys
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<400> 143
Cys Cys Glu Val Cys Cys Asn Pro Ala Cys Thr Gly Cys
<210> 144
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<222> 8
<223> Xaa = Glu
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<223> Xaa = Leu, Ile, Lys, Arg, Trp, Tyr or Phe
<220>
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<222> 12
<223> Xaa = Asn
<220>
<221>VARIANT
<222> 13
<223> Xaa = Pro
<220>
<221>VARIANT
<222> 14
<223> Xaa = Ala
<220>
<221>VARIANT
<222> 16
<223> Xaa = Thr, Ala, Lys, Arg, Trp
<220>
<221>VARIANT
<222> 17
<223> Xaa = Gly
<220>
<221>VARIANT
<222> 19
<223> Xaa = Tyr or Leu
<220>
<221>VARIANT
<222> 20, 21
<223> Xaa20 = Asp; Xaa21 = Phe; or missing
Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Cys Cys Xaa Xaa Xaa Cys Xaa
Xaa Cys Xaa Xaa Xaa
<210> 145
<211> 21
<212> PRT
<213> Artificial Sequence
<220>
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<220>
<221>VARIANT
<222> 1, 2, 3, 4, 5
<223> Xaa1= Asn, Xaa2 = Ser, Xaa3 = Ser, Xaa4 = Asn,
      Xaa5 = Tyr; or mising
<220>
<221>VARIANT
<222> 1, 2, 3, 4
<223> Xaa = missing
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<222> 5
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<220>
<221>VARIANT
<222> 8
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<220>
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<223> Xaa = Leu, Ile, Val, Ala, Lys, Arg, Trp, Tyr or
      Phe
<220>
<221>VARIANT
<222> 12
<223> Xaa = Asn, Tyr, Asp or Ala
<220>
<221>VARIANT
<222> 13
<223> Xaa = Pro or Gly
<220>
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<223> Xaa = Ala, Leu, Ser, Gly, Val, Glu, Gln, Ile, Leu,
      Lys, Arg, and Asp
<220>
<221>VARIANT
<222> 16
<223> Xaa = Thr, Ala, Asn, Lys, Arg
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<221>VARIANT
<222> 17
<223> Xaa = Gly, Pro or Ala
<220>
<221>VARIANT
<222> 19
<223> Xaa = Trp, Tyr, Phe or Leu
<220>
<221>VARIANT
<222> 19-21
<223> Xaa = Asp, Phe or missing; or Xaa20 =
      Asn, or Glu and Xaa21 is missing; or Xaa19, Xaa20,
      Xaa21 = is missing
<400> 145
Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Cys Cys Xaa Xaa Xaa Cys Xaa
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1
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Xaa Cys Xaa Xaa Xaa
<210> 146
<211> 21
<212> PRT
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<223> Synthetically generated peptide
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<221>VARIANT
<222> 1, 2, 3, 4, 5
<223> Xaa = missing
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<221>VARIANT
<222> 8
<223> Xaa = Glu
<220>
<221>VARIANT
<222> 9
<223> Xaa = Leu, Ile, Lys, Arg, Trp, Tyr, or Phe
<220>
<221>VARIANT
<222> 12
<223> Xaa = Asn
<220>
<221>VARIANT
<222> 13
<223> Xaa = Pro
<220>
<221>VARIANT
<222> 14
<223> Xaa = Ala
<220>
<221>VARIANT
<222> 16
<223> Xaa = Thr, Ala, Lys, Arg, Trp or X16 = any amino
      acid; or X16 = Thr, Ala, Lys, Arg, Trp or any
      non-aromatic amino acid
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<221>VARIANT
<222> 17
<223> Xaa = Gly
<220>
<221>VARIANT
<222> 19
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<223> Xaa = Tyr or Leu
<220>
<221>VARIANT
<222> 20, 21
<223> Xaa20 = Asp, Xaa21 = Phe or missing
<400> 146
Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Cys Cys Xaa Xaa Cys Xaa
                                    10
Xaa Cys Xaa Xaa Xaa
            20
<210> 147
<211> 21
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<221>VARIANT
<222> 1, 2, 3, 4, 5
<223> Xaa = Xaa1 = Asn, Xaa2 = Ser, Xaa3 = Ser, Xaa4 =
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      mising and Xaa5 = Asn, Trp, Tyr, Asp, Ile, Thr
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<223> Xaa = Leu, Ile, Val, Ala, Lys, Arg, Trp, Tyr or
      Phe
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<221>VARIANT
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<220>
<221>VARIANT
<222> 13
<223> Xaa = Pro or Gly
<220>
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<222> 14
<223> Xaa = Ala, Leu, Ser, Gly, Val, Glu, Gln, Ile, Leu,
      Lys, Arg or Asp
<220>
<221>VARIANT
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<222> 16
<223> Xaa = Thr, Ala, Asn, Lys, Arg, Trp
<220>
<221>VARIANT
<222> 17
<223> Xaa = Gly, Pro or Ala
<220>
<221>VARIANT
<222> 19
<223> Xaa = Trp, Tyr, Phe or Leu; or Xaa = Lys or Arg
<220>
<221>VARIANT
<222> 20
<223> Xaa = Asp, Phe or missing; or
      Xaa20 = Asn or Glu and Xaa21 is missing
<220>
<221>VARIANT
<222> 19, 21
<223> Xaa is miising
<400> 147
Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Cys Cys Xaa Xaa Cys Xaa
                                    10
Xaa Cys Xaa Xaa Xaa
            20
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<222> 1, 2, 3, 4, 5
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<223> Xaa = Glu
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<222> 9
<223> Xaa = Leu, Ile, Lys, Arg, Trp, Tyr or Phe
<220>
<221>VARIANT
<222> 12
<223> Xaa = Asn
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<223> Xaa = Pro
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<221>VARIANT
<222> 14
<223> Xaa = Ala
<220>
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<221>VARIANT
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<223> Xaa = Gly
<220>
<221>VARIANT
<222> 19
<223> Xaa = Tyr or Leu; or Xaa = Lys or Arg
<220>
<221>VARIANT
<222> 20, 21
<223> Xaa = Asp, Phe or is missing
<400> 148
Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Cys Cys Xaa Xaa Xaa Cys Xaa
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                                   10
Xaa Cys Xaa Xaa Xaa
            20
<210> 149
<211> 21
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<222> 1, 2, 3, 4
<223> Xaa = is missing
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<221>VARIANT
<222> 5
<223> Xaa = Asn
<220>
<221>VARIANT
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<222> 9
<223> Xaa = Trp, Tyr or Phe;
<220>
<221>VARIANT
<222> 16
<223> Xaa = Thr or Ala
<220>
<221>VARIANT
<222> 19
<223> Xaa = Trp, Tyr, Phe
<220>
<221>VARIANT
<222> 20, 21
<223> Xaa = Asp, Phe
<400> 149
Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Cys Cys Asn Pro Ala Cys Xaa
                                    10
Gly Cys Xaa Xaa Xaa
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